

HT1220 Report

RECORD INFORMATION

 Gene ID:
 1220

 Sequence ID:
 1220

 Protein ID:
 1220

Sequence name: thrombospondin 1, alt. transcript 1

Genome: nucleus
Taxon: Homo sapiens

Locus: 1220

Common Name: thrombospondin 1

Role ID: 40

Coding sequence length: Transcript sequence length:

Transcript sequence length: Expression data:

3513 nt 5722 nt THC201673

ACCESSION DATA

HT1220 is derived from accessions(s):

SP:P07996 (THROMBOSPONDIN 1 PRECURSOR.)
GB:X04665 (Human mRNA for thrombospondin)
GB:X14787 (Human mRNA for thrombospondin)
GB:U12471 (thrombospondin-p50 {Homo sapiens})
GB:M99425 (Human thrombospondin mRNA, 3' end.)
PIR:G01478 (thrombospondin-p50 - human (fragment))
GB:U12471 (Human thrombospondin-1 gene, partial cds.)
GB:J04835 (Human thrombospondin gene, exons 1, 2 and 3.)

GB:M25631 (Homo sapiens (clone lambda-TS-33) thrombospondin (THBS) mRNA, 5' end.)

ALTERNATIVE SPLICE INFORMATION

Alternative splice forms for this gene:

HT3987 thrombospondin 1, alt. transcript 2

MAPPING DATA

GDB accession(s) for this gene:

GDB ID: Symbol

gdb:120438 THBS1

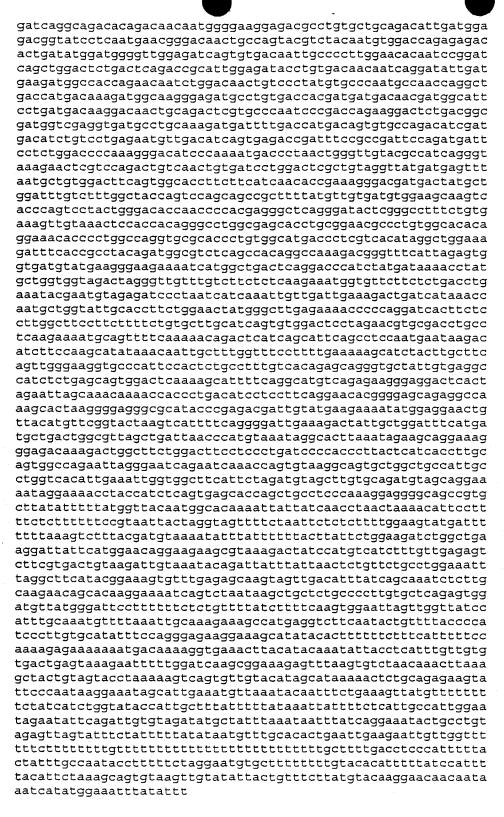
cDNA FEATURES

Feature	End 5	End 3
coding_seq 3'UT spjunc_h	112 3625 1235	3624 5722 1236

SEQUENCE

nucleotide:

ggacgcacaggcattccccgcgcccctccagccctcgccgccctcgccaccgctcccggc cgccgcgctccggtacacacaggatccctgctgggcaccaacagctccaccatggggctg tetqqqeqeeqaetqqtqaaqqqeeeqaeeetteeaqeeeaqettteeqeateqaqqat gccaacctgatccccctgtgcctgatgacaagttccaagacctggtggatgctgtgcgg gcagaaaagggtttcctccttctggcatccctgaggcagatgaagaagacccggggcacg ctgctggccctggagcggaaagaccactctggccaggtcttcagcgtggtgtccaatggc aaggegggcaeeetggaeeteageetgaeegteeaaggaaageageaegtggtgtetgtg gaagaagctctcctggcaaccggccagtggaagagcatcaccctgtttgtgcaggaagac agggcccagctgtacatcgactgtgaaaagatggagaatgctgagttggacgtccccatc caaagcgtcttcaccagagacctggccagcatcgccagactccgcatcgcaaaggggggc gtcaatgacaatttccagggggtgctgcagaatgtgaggtttgtctttggaaccacacca gaagacatcctcaggaacaaaggctgctccagctctaccagtgtcctcctcacccttgac aaggacttgcaagccatctgcggcatctcctgtgatgagctgtccagcatggtcctggaa ctcaggggcctgcgcaccattgtgaccacgctgcaggacagcatccgcaaagtgactgaa gagaacaaagagttggccaatgagctgaggcggcctcccctatgctatcacaacggagtt cagtacagaaataacgaggaatggactgttgatagctgcactgagtgtcactgtcagaac tcagttaccatctgcaaaaaggtgtcctgccccatcatgccctgctccaatgccacagtt cctgatggagaatgctgtcctcgctgttggcccagcgactctgcggacgatggctgtct ccatggtccgagtggacctcctgttctacgagctgtggcaatggaattcagcagcgcggc cgctcctgcgatagcctcaacaaccgatgtgagggctcctcggtccagacacggacctgc cacattcaggagtgtgacaaaagatttaaacaggatggtggctggagccactggtccccg tggtcatcttgttctgtgacatgtggtgatggtgatcacaaggatccggctctgcaac tctcccagcccccagatgaatgggaaaccctgtgaaggcgaagcgggggagaccaaagcc tgcaagaaagacgcctgccccatcaatggaggctggggtccttggtcaccatgggacatc tgttctgtcacctgtggaggaggggtacagaaacgtagtcgtctctgcaacaaccccgca ccccagtttggaggcaaggactgcgttggtgatgtaacagaaaccagatctgcaacaag caggactgtccaattgatggatgcctgtccaatccctgctttgccggcgtgaagtgtact agctaccctgatggcagctggaaatgtggtgcttgtccccctggttacagtggaaatggc atccagtgcacagatgttgatgagtgcaaagaagtgcctgatgcctgcttcaaccacaat ttcaccggctcacagcccttcggccagggtgtcgaacatgccacggccaacaacaggtg tgcaagccccgtaacccctgcacggatgggacccacgactgcaacaagaacgccaagtgc aactacctgggccactatagcgaccccatgtaccgctgcgagtgcaagcctggctacgct gtgtgcgtggccaatgcgacttaccactgcaaaaaggataattgccccaaccttcccaac $\verb|tcagggcaggaagactatgacaaggatggaattggtgatgcctgtgatgatgacgatgac|$ ${\tt aatgataaaattccagatgacagggacaactgtccattccattacaacccagctcagtat}$ gactatgacagagatgatgtgggagaccgctgtgacaactgtccctacaaccacaaccca



protein:

MGLAWGLGVLFLMHVCGTNRIPESGGDNSVFDIFELTGAARKGSGRRLVKGPDPSSPAFR

IEDANLIPPVPDDKFQDLVDAVRAEKGFLLLASLRQMKKTRGTLLALERKDHSGQVFSVV SNGKAGTLDLSLTVQGKQHVVSVEEALLATGQWKSITLFVQEDRAQLYIDCEKMENAELD VPIQSVFTRDLASIARLRIAKGGVNDNFQGVLQNVRFVFGTTPEDILRNKGCSSSTSVLL TLDNNVVNGSSPAIRTNYIGHKTKDLQAICGISCDELSSMVLELRGLRTIVTTLQDSIRK VTEENKELANELRRPPLCYHNGVQYRNNEEWTVDSCTECHCQNSVTICKKVSCPIMPCSN ATVPDGECCPRCWPSDSADDGWSPWSEWTSCSTSCGNGIQQRGRSCDSLNNRCEGSSVQT RTCHIQECDKRFKQDGGWSHWSPWSSCSVTCGDGVITRIRLCNSPSPQMNGKPCEGEARE TKACKKDACPINGGWGPWSPWDICSVTCGGGVQKRSRLCNNPAPQFGGKDCVGDVTENQI ${\tt CNKQDCPIDGCLSNPCFAGVKCTSYPDGSWKCGACPPGYSGNGIQCTDVDECKEVPDACF}$ NHNGEHRCENTDPGYNCLPCPPRFTGSQPFGQGVEHATANKQVCKPRNPCTDGTHDCNKN AKCNYLGHYSDPMYRCECKPGYAGNGIICGEDTDLDGWPNENLVCVANATYHCKKDNCPN LPNSGQEDYDKDGIGDACDDDDDDKIPDDRDNCPFHYNPAQYDYDRDDVGDRCDNCPYN HNPDQADTDNNGEGDACAADIDGDGILNERDNCQYVYNVDQRDTDMDGVGDQCDNCPLEH NPDQLDSDSDRIGDTCDNNQDIDEDGHQNNLDNCPYVPNANQADHDKDGKGDACDHDDDN DGIPDDKDNCRLVPNPDQKDSDGDGRGDACKDDFDHDSVPDIDDICPENVDISETDFRRF QMIPLDPKGTSQNDPNWVVRHQGKELVQTVNCDPGLAVGYDEFNAVDFSGTFFINTERDD DYAGFVFGYQSSSRFYVVMWKQVTQSYWDTNPTRAQGYSGLSVKVVNSTTGPGEHLRNAL WHTGNTPGQVRTLWHDPRHIGWKDFTAYRWRLSHRPKTGFIRVVMYEGKKIMADSGPIYD KTYAGGRLGLFVFSQEMVFFSDLKYECRDP







HT2143 Report

RECORD INFORMATION

 Gene ID:
 2081

 Sequence ID:
 2143

 Protein ID:
 2125

Sequence name: thrombospondin 4

Genome: nucleus
Taxon: Homo sapiens

Locus: 2081

Common Name: thrombospondin 4

Role ID:

Coding sequence length: Transcript sequence length:

Expression data:

2886 nt 3074 nt THC168897

ACCESSION DATA

HT2143 is derived from accessions(s):

SP:P35443 (THROMBOSPONDIN 4 PRECURSOR.)
GB:Z19585(thrombospondin-4 {Homo sapiens})
GB:Z19585(H.sapiens mRNA for thrombospondin-4)
PIR:A55710 (thrombospondin 4 precursor - human)

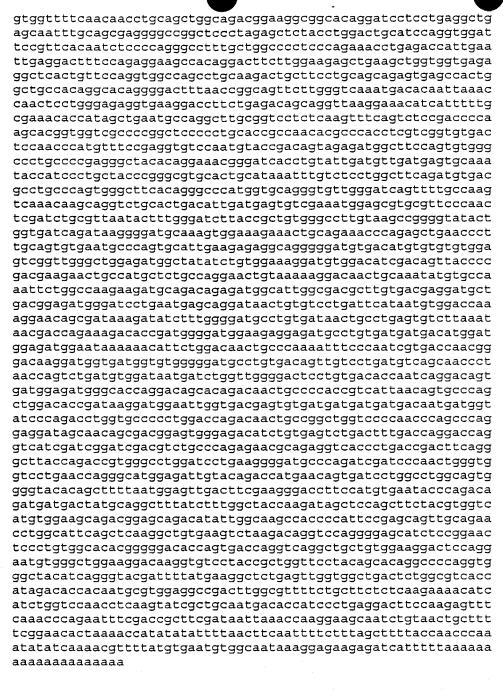
cDNA FEATURES

Feature	End 5	End 3
		.
coding_seq	28	2913
3'UT	2914	3074

SEQUENCE

nucleotide:

gaattccggggagcaggaagagccaacatgctggccccgcgcgggagccgccgtcctcctg ctgcacctggtcctgcagcggtggctagcggcaggcgcccaggccaccccccaggtcttt gaccttctcccatcttccagtcagaggctaaacccaggcgctctgctgccagtcctgaca gaccccgccctgaatgatctctatgtgatttccaccttcaagctgcagactaaaagttca gccaccatcttcggtctttactcttcaactgacaacagtaaatattttgaatttactgtg atgqgacgcttaagcaaagccatcctccgttacctgaagaacgatggaaggtgcattt



protein:

MLAPRGAAVLLLHLVLQRWLAAGAQATPQVFDLLPSSSQRLNPGALLPVLTDPALNDLYV ISTFKLQTKSSATIFGLYSSTDNSKYFEFTVMGRLSKAILRYLKNDGKVHLVVFNNLQLA DGRRHRILLRLSNLQRGAGSLELYLDCIQVDSVHNLPRAFAGPSQKPETIELRTFQRKPQ DFLEELKLVVRGSLFQVASLQDCFLQQSEPLAATGTGDFNRQFLGQMTQLNQLLGEVKDL LRQQVKETSFLRNTIAECQACGPLKFQSPTPSTVVAPAPPAPPTRPPRRCDSNPCFRGVQ CTDSRDGFQCGPCPEGYTGNGITCIDVDECKYHPCYPGVHCINLSPGFRCDACPVGFTGP MVQGVGISFAKSNKQVCTDIDECRNGACVPNSICVNTLGSYRCGPCKPGYTGDQIRGCKV ERNCRNPELNPCSVNAQCIEERQGDVTCVCGVGWAGDGYICGKDVDIDSYPDEELPCSAR NCKKDNCKYVPNSGQEDADRDGIGDACDEDADGDGILNEQDNCVLIHNVDQRNSDKDIFG DACDNCLSVLNNDQKDTDGDGRGDACDDDMDGDGIKNILDNCPKFPNRDQRDKDGDGVGD

<u>-</u>

ACDSCPDVSNPNQSDVDNDLVGDSCDTNQDSDGDGHQDSTDNCPTVINSAQLDTDKDGIG
DECDDDDDDGIPDLVPPGPDNCRLVPNPAQEDSNSDGVGDICESDFDQDQVIDRIDVCP
ENAEVTLTDFRAYQTVGLDPEGDAQIDPNWVVLNQGMEIVQTMNSDPGLAVGYTAFNGVD
FEGTFHVNTQTDDDYAGFIFGYQDSSSFYVVMWKQTEQTYWQATPFRAVAEPGIQLKAVK
SKTGPGEHLRNSLWHTGDTSDQVRLLWKDSRNVGWKDKVSYRWFLQHRPQVGYIRVRFYE
GSELVADSGVTIDTTMRGGRLGVFCFSQENIIWSNLKYRCNDTIPEDFQEFQTQNFDRFD





Rol AA AII AA	ss Z	A	
ANNT	O	೦	
. Mutation T. Ref NT	Missense A	9	
, Flanking Seq	TOGATOGCTGGCCCAINGITGAGAACCTGGTGTG	354, Informbospondin 4 GAGTGTCGAAATBGAGGCCGTGCGTTCCCAACT	
e Description	2210 THBS1, thrombospondin 1	1186 THBS4, Informbosporiditi	
Sequence ID	HIT1471220 mRNA	3385u2 HTHT2143 mRNA	
Poly ID	6334114	3355112	

Figure 3